In the Claims:

- 1-39. (Canceled)
- 40. (Currently amended) A method comprising:
- a) obtaining a target protein comprising a -SH group, masked -SH group, or activated -SH group;
- b) combining said target protein, in aqueous solution, with a library simultaneously containing at least two non-oligomeric ligand candidates wherein said <u>non-oligomeric</u> ligand candidates each comprise a disulfide bond, and wherein said <u>non-oligomeric</u> ligand candidates each are less than about 2000 daltons in size, under disulfide-exchange conditions, in the presence of a reducing agent, wherein at least one <u>non-oligomeric</u> ligand candidate binds to the target protein and forms a disulfide bond with the target protein to yield a target protein-ligand conjugate; and
- c) determining the identity of the non-oligomeric ligand present in said target proteinligand conjugate by subjecting said conjugate directly, without prior fragmentation and without liberation of the ligand from said conjugate, to mass spectrometry analysis.
- 41. (Previously presented) The method of claim 40 wherein the ligand is less than 1500 daltons.
 - 42. (Canceled)
- 43. (Previously presented) The method of claim 40 wherein the ligand is less than 750 daltons.
 - 44-46. (Canceled)
- 47. (Currently amended) The method of claim <u>40</u> 45 or claim 46 wherein the reducing agent is 2-mercaptoethanol.
 - 48-58. (Canceled)
- 59. (Currently amended) A method for identifying a non-oligomeric ligand that binds to a target protein wherein said ligand less than about 750 2000 daltons in size, said method comprising:
- a) obtaining said target protein comprising a -SH group, masked -SH group, or activated -SH group;

- b) combining said target protein, in aqueous solution, with a library containing at least two non-oligomeric ligand candidates in a mixture wherein said <u>non-oligomeric</u> ligand candidates each comprise a disulfide bond, and wherein said <u>non-oligomeric</u> ligand candidates are each less than about 750 2000 daltons in size, under disulfide exchange conditions, in the presence of a reducing agent, wherein at least one <u>non-oligomeric</u> ligand candidate binds to the target protein and forms a covalent disulfide bond with the target protein to yield a <u>covalent</u> target protein-ligand conjugate;
 - (c) separating the covalent target protein-ligand conjugate from the mixture; and
- (d) determining the identity of the ligand present in said conjugate <u>by subjecting said</u> conjugate directly, without prior fragmentation and without liberation of the ligand from said conjugate, to mass spectrometry analysis.

60-63. (Canceled)

- 64. (Previously presented) The method of claim 40 wherein said -SH group, masked SH group, or activated -SH group is associated with part of a cysteine residue of said target protein.
 - 65. (New) The method of claim 59 wherein the ligand is less than 1500 daltons.
 - 66. (New) The method of claim 59 wherein the ligand is less than 750 daltons.
 - 67. (New) The method of claim 59 wherein the reducing agent is 2-mercaptoethanol.